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RAW SEQUENCE LISTING

DATE: 08/30/2001

PATENT APPLICATION: US/09/701,536

TIME: 07:54:47

Input Set : A:\W098706.txt

Output Set: N:\CRF3\08302001\1701536.raw

## ENTERED

```
SEQUENCE LISTING
      4 (1) GENERAL INFORMATION:
             (i) APPLICANT: Chang, Gwong-Jen J
      6
      8
            (ii) TITLE OF INVENTION: Nucleic Acid Vaccines for
      9
                                      Prevention of Flavivirus Infection
     11
           (iii) NUMBER OF SEQUENCES: 6
     -13.
           _(iv) CORRESPONDENCE ADDRESS:
                  (A) ADDRESSEE: Needle & Rosenberg, P.C.
     14
     15
                  (B) STREET: 127 Peachtree St., NE., Ste. 1200, The Candler Bldg.
     16
                  (C) CITY: Atlanta
    二17
                  (D) STATE: GA
    18
                  (E) COUNTRY: USA
                  (F) ZIP: 30303-1811
             (V) COMPUTER READABLE FORM:
                  (A) MEDIUM TYPE: Diskette
                  (B) COMPUTER: IBM Compatible
                  (C) OPERATING SYSTEM: Windows
                  (D) SOFTWARE: FastSEQ for Windows Version 4.0
   Ū27
            (vi) CURRENT APPLICATION DATA:
                  (A) APPLICATION NUMBER: US/09/701,536
C--> 28
C-->_29
                  (B) FILING DATE: 29-Nov-2000
                  (C) CLASSIFICATION: U.S. National Phase
   730
   32ليا
           (vii) PRIOR APPLICATION DATA:
                  (A) APPLICATION NUMBER: PCT/US99/12298
                  (B) FILING DATE: June 3, 1999
          (viii) ATTORNEY/AGENT INFORMATION:
   37
                  (A) NAME: Miller, Mary L.
     38
                  (B) REGISTRATION NUMBER: 39,303
     39
                  (C) REFERENCE/DOCKET NUMBER: 14014.0332U2
            (ix) TELECOMMUNICATION INFORMATION:
     41
     42
                  (A) TELEPHONE: 404-688-0770
                  (B) TELEFAX: 404-688-9880 ·
     43
     44
                  (C) TELEX:
        (2) INFORMATION FOR SEQ ID NO: 1:
     47
             (i) SEQUENCE CHARACTERISTICS:
     49
     50
                  (A) LENGTH: 48 base pairs
     51
                  (B) TYPE: nucleic acid
     52
                  (C) STRANDEDNESS: single
     53
                  (D) TOPOLOGY: linear
     55
            (ii) MOLECULE TYPE: Other
     56
            (ix) FEATURE:
     63
                  (A) NAME/KEY: Other
     59
                  (B) LOCATION: 25...48
     60
                  (D) OTHER INFORMATION:
     68
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
    . 70
        CTTGGTACCT CTAGAGCCGC CGCC ATG GGC AGA AAG CAA AAC AAA AGA
                                                                                  48
     71
                                     Met Gly Arg Lys Gln Asn Lys Arg
```



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72
 75 (2) INFORMATION FOR SEQ ID NO: 2:
 77
          (i) SEQUENCE CHARACTERISTICS:
 78
               (A) LENGTH: 50 base pairs
 79
               (B) TYPE: nucleic acid
 80
              (C) STRANDEDNESS: single
 81
               (D) TOPOLOGY: linear
         (ii) MOLECULE TYPE: Other
 83
 84
         (ix) FEATURE:
 86
               (A) NAME/KEY: Other
               (B) LOCATION: 1...50
 87
               (D) OTHER INFORMATION: Amplimer c14DV2453
 88
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
 90
 92
     TTTTCTTTG CGCCGCTCA AACTTAAGCA TGCACATTGG TCGCTAAGAA
 94 (2) INFORMATION FOR SEQ ID NO: 3: -------
□96
          (i) SEQUENCE CHARACTERISTICS:
⊈97
               (A) LENGTH: 48 base pairs
98
               (B) TYPE: nucleic acid
99
               (C) STRANDEDNESS: single
100
                (D) TOPOLOGY: linear
102
          (ii) MOLECULE TYPE: Other
103
         (ix) FEATURE:
[110
                (A) NAME/KEY: Other
 106
                (B) LOCATION: 25...48
 107
                (D) OTHER INFORMATION:
115
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
<sup>1</sup>117
      CTTGGTACCT CTAGAGCCGC CGCC ATG CGT TCC CAT GAT GTT CTG ACT
                                                                               48
118
                                  Met Arg Ser His Asp Val Leu Thr
119
                                                    5
122 (2) INFORMATION FOR SEQ ID NO: 4:
124
           (i) SEQUENCE CHARACTERISTICS:
 125
                (A) LENGTH: 41 base pairs
 126
                (B) TYPE: nucleic acid
 127
                (C) STRANDEDNESS: single
                (D) TOPOLOGY: linear
 128
 130
         (ii) MOLECULE TYPE: Other
 131
         (ix) FEATURE:
 133
                (A) NAME/KEY: Other
 134
                (B) LOCATION: 1...41
 135
                (D) OTHER INFORMATION: Amplimer cYFDV2452
 138
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
      TTTTCTTTTG CGGCCGCTCA CGCCCCAACT CCTAGAGAAA C
                                                                               41
 142 (2) INFORMATION FOR SEQ ID NO: 5:
 144
          (i) SEQUENCE CHARACTERISTICS:
 145
                (A) LENGTH: 54 base pairs
 146
                (B) TYPE: nucleic acid
                (C) STRANDEDNESS: single
 147
 148
                (D) TOPOLOGY: linear
 150
         (ii) MOLECULE TYPE: Other
```





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151	(ix) FEATURE:	
158	(A) NAME/KEY: Other	
· 154	(B) LOCATION: 2554	
155	(D) OTHER INFORMATION:	
163	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:	
165	CTTGGTACCT CTAGAGCCGC CGCC ATG TCT AAA AAA AGA GGA GGG ACC AGA	51
166	Met Ser Lys Lys Arg Gly Gly Thr Arg	
167	1 5	
169	TCG	54
170	Ser	
171	10	
174	(2) INFORMATION FOR SEQ ID NO: 6:	
1.7.6	· · · · · · · · · · · · ·	
177	(A) LENGTH: 38-base pairs	
178	(B) TYPE: nucleic acid	
<b>[</b> 179	(C) STRANDEDNESS: single	
<b>_</b> 180	(D) TOPOLOGY: linear	
182	(ii) MOLECULE TYPE: Other	
<b>=</b> 183	(ix) FEATURE:	
<u></u> 185	(A) NAME/KEY: Other	
186 187	(B) LOCATION: 138	
¥ 187	(D) OTHER INFORMATION: Amplimer cSLEDV2449	
190	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:	
192	TTTTCTTTTG CGGCCGCTTA GGCTTGCACG CTGGTTGC	38
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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/701,536

DATE: 08/30/2001 TIME: 07:54:48

Input Set : A:\W098706.txt

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L:28 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:] L:29 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]

